Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1. (Currently Amended) A method of producing an immunoglobulin constant region, comprising:

transforming a prokaryotic cell with a recombinant expression vector including a nucleotide sequence encoding an <u>signal sequence</u> isolated from *E. coli*—derived signal sequence and a nucleotide sequence encoding an immunoglobulin constant region;

culturing a resulting transformant; and

isolating and purifying the immunoglobulin constant region expressed by the transformant,

wherein the signal sequence is a heat-stable enterotoxin II signal sequence.

- 2. (Original) The method according to claim 1, wherein the immunoglobulin constant region is selected from the group consisting of constant regions from IgG, IgA, IgM, IgE, IgD, and combinations and hybrids thereof.
- 3. (Original) The method according to claim 2, wherein the IgG is selected from the group consisting of constant regions from IgG1, IgG2, IgG3, IgG4, and combinations and hybrids thereof.
- 4. (Original) The method according to claim 3, wherein the immunoglobulin constant region is an IgG4 constant region.
- 5. (Original) The method according to claim 4, wherein the immunoglobulin constant region is a human aglycosylated IgG4 constant region.

- 6. (Currently Amended) The method according to claim 1, wherein the immunoglobulin constant region is composed of one to four domains selected from the group[[s]] consisting of C_H1, C_H2, C_H3, and C_H4 and C_L domains.
- 7. (Original) The method according to claim 6, wherein the immunoglobulin constant region further comprises a hinge region.
- 8. (Original) The method according to claim 1, wherein the recombinant expression vector comprises a nucleotide sequence encoding a heavy chain constant region and a nucleotide sequence encoding a light chain constant region.
- 9. (Original) The method according to claim 1, wherein the immunoglobulin constant region has an amino acid sequence represented by SEQ ID NO. 21, 22, 23, 24, 25, 27, 29, 30, 34 or 35.

10. (Canceled)

- 11. (Original) The method according to claim 10, wherein the heat-stable enterotoxin II signal peptide has an amino acid sequence represented by SEQ ID NO. 36, 37, 38, 39, 40, 41, 42, 43, 44, 45 or 46.
- 12. (Currently Amended) The method according to claim 1, wherein the recombinant expression vector—is—pSTIIGICH1_3, pSTIIGCG1Fe, pSTIIGCG1SFe, pSTIIGCG1SFe,

- 13. (Currently Amended) The method according to claim 1, wherein the transformant is selected from the group consisting of E. coli BL21/pSTIIGICH1_3(HM10935; Deposit No. KCCM-10600), BL21/pSTIIdCG1Fc (HM10927; Deposit No. KCCM-10588), BL21/pSTIIdCG1SFc (HM10928; Deposit No. KCCM-10589), BL21/pSTIIdCG1SFc (HM10929; Deposit No. KCCM-10594), BL21/pSTIIG1Mo (HM10930; Deposit No. KCCM-10595), BL21/pSTIIdCG2Fc (HM10936), BL21/pSTIIdCG4Fc (HM10932; Deposit No. KCCM-10597), BL21/pSTIIGICH1_3_(HM10931; Deposit No. KCCM-10596), BL21/pSTIIG4Mo (HM10933; Deposit No. KCCM-10598), or and BL21/pSTIIG4H_K (HM10934; Deposit No. KCCM-10599).
- 14. (Original) The method according to claim 1, wherein the transformant is *E. coli*.
- 15. (Withdrawn) An immunoglobulin constant region prepared by the method of claim 1.
- 16. (New) The method according to claim 1, wherein the immunoglobulin constant region comprises a CL domain or one to four domains selected from the group consisting of C_H1, C_H2, C_H3, and C_H4 domains.

Amendments to the Drawings:

The attached drawing sheet includes a change to Figure 4. This sheet replaces the original sheet Fig. 4.

Attachment: Replacement Sheet